RAW SEQUENCE LISTING PATENT APPLICATION US/08/013,413A

DATE: 09/17/93 TIME: 15:21:51



1		SEQUENCE LISTING	
2 3	(1) Ge	eneral Information:	
4			ENTE
5 6	(i)	APPLICANT: Le, Junming	— 10 U C
7		Vilcek, Jan Daddona, Peter E.	
8		Ghrayeb, John	
9		Knight, David M.	
10		Siegel, Scott A.	
11			
12	(ii)	TITLE OF INVENTION: MONOCLONAL AND CHIMERIC	ANTIBODIES '
13		SPECIFIC FOR HUMAN TUMOR NECROSIS FACTOR	
14 15	(:::)	NUMBER OF CECHENCES. F	
16	(111)	NUMBER OF SEQUENCES: 5	(6,7.20
17	(iv)	CORRESPONDENCE ADDRESS:	1600
18	(= - ,	(A) ADDRESSEE: Browdy and Neimark	\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \
19		(B) STREET: 419 Seventh Street, N.W.	·
20		(C) CITY: Washington	
21		(D) STATE: D.C.	
22		(E) COUNTRY: USA	
23 24		(F) ZIP: 20004	
24 25	(37)	COMPUTER READABLE FORM:	
26	(•)	(A) MEDIUM TYPE: Floppy disk	
27		(B) COMPUTER: IBM PC compatible	
28		(C) OPERATING SYSTEM: PC-DOS/MS-DOS	
29		(D) SOFTWARE: PatentIn Release #1.0,	
30	Ver	rsion #1.25	
31	()	OUDDOWN ADDITIONAL DAMA	
32 33	(V1)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: 08/013,413	
34		(B) FILING DATE: 02-FEB-1993	
35		(C) CLASSIFICATION: 436	
36		, , , , , , , , , , , , , , , , , , , ,	
37	(vii)	PRIOR APPLICATION DATA:	
38		(A) APPLICATION NUMBER: US 07/943,852	
39		(B) FILING DATE: 11-SEP-1992	
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41 42	(V11)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: US 07/853,606	
43		(B) FILING DATE: 18-MAR-1992	
44		(b) Italia biil. 10 ital 1991	
45	(vii)	PRIOR APPLICATION DATA:	
46		(A) APPLICATION NUMBER: US 07/670,827	
47		(B) FILING DATE: 18-MAR-1991	
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49	(1X)	TELECOMMUNICATION INFORMATION:	•
50 51		(A) TELEPHONE: 202-628-5197 (B) TELEFAX: 202-737-3528	
ÜΙ		(D) IBBERA. 202-131-3526	

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65 66			Arg	Ser	Ser		Arg	Thr	Pro	Ser	_	Lys	Pro	Val	Ala		Val
66 67		1				5					10					15	
68		Val	Ala	Asn	Pro	Gln	Ala	Glu	Glv	Gln	Leu	Gln	Trp	Leu	Asn	Ara	Ara
69					20				1	25			1-		30	5	5
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71		Ala	Asn		Leu	Leu	Ala	Asn	_	Val	Glu	Leu	Arg	-	Asn	Gln	Leu
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77		Lys	Gly	Gln	Gly	Cys	Pro	Ser	Thr	His	Val	Leu	Leu	Thr	His	Thr	Ile
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83		Ile	Lys	Ser	Pro	Cys	Gln	Arg	Glu	Thr	Pro	Glu	Gly	Ala	Glu	Ala	Lys
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86		Pro	Trp	_	GLu	Pro	IIe	Tyr		GLY	GLY	Val	Phe		Leu	GIu	Lys
87 88				115					120					125			
89		Gly	Asp	Arq	Leu	Ser	Ala	Glu	Ile	Asn	Arq	Pro	Asp	Tyr	Leu	qaA	Phe
90		•	130					135					140	•		-	
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92			Glu	Ser	Gly	Gln		Tyr	Phe	Gly	Ile		Ala	Leu			
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114	GAC	ATC	TTG	CTG	ACT	CAG	TCT	CCA	GCC	ATC	CTG	TCT	GTG	AGT	CCA	GGA	48
115	Asp	Ile	Leu	Leu	Thr	Gln	Ser	Pro	Ala	Ile	Leu	Ser	Val	Ser	Pro	Gly	
116	ī				5					10					15	-	
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118	GAA	AGA	GTC	AGT	ттс	TCC	TGC	AGG	GCC	ΔСΤ	CAG	ጥጥር	СТТ	GGC	тса	AGC	96
119				Ser													70
120	Gru	Arg	vai		FIIC	Ser	Cys	ALG		261	GIII	FILE	vai	_	261	261	
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123	Ile	His	_	Tyr	GIn	GIn	Arg		Asn	GТУ	Ser	Pro	Arg	Leu	Leu	Ile	
124			35					40					45				
125																	
126	AAG	TAT	GCT	TCT	GAG	TCT	ATG	TCT	GGG	ATC	CCT	TCC	AGG	TTT	AGT	GGC	192
127	Lys	Tyr	Ala	Ser	Glu	Ser	Met	Ser	Gly	Ile	Pro	Ser	Arg	Phe	Ser	Gly	
128		50					55		_			60	_			_	
129																	
130	AGT	GGA	TCA	GGG	ACA	GAT	TTT	ACT	CTT	AGC	ATC	AAC	ACT	GTG	GAG	TCT	240
131				Gly													
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				GCA													288
135	GIU	Asp	TTE	Ala	_	Tyr	Tyr	Cys	GIN		Ser	His	Ser	Trp		Pne	
136					85					90					95		
137																	
138	ACG	TTC	GGC	TCG	GGG	ACA	TAA	TTG	GAA	GTA	AAA						321
139	Thr	Phe	Gly	Ser	Gly	Thr	Asn	Leu	Glu	Val	Lys						
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154 155	Asp 1	Ile	Leu	Leu	Thr 5	Gln	Ser	Pro	Ala	Ile 10	Leu	Ser	Val	Ser	Pro 15	Gly	
156																	
157	Glu	λνα	17a l	Ser	Dhe	202	Cara	7/ ***	777	202	Cln	Dho	3707	C1	000	Com	
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160	Ile	His	Trp	Tyr	Gln	Gln	Arq	Thr	Asn	Gly	Ser	Pro	Arq	Leu	Leu	Ile	
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166	Ser	Glv	Ser	Gly	Thr	asp	Phe	Thr	Leu	Ser	Ile	Asn	Thr	Val	Glu	Ser	
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172	Thr	Phe	Glv	Ser	Glv	Thr	Asn	Leu	Glu	Val	Lvs						
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193				CTT													48
194	Glu	Val	Lys	Leu	GLu	GLu	Ser	GГУ	Gly	GГУ	Leu	Val	GIn	Pro	GГУ	GГУ	
195	1				5					10					15		
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197	TCC	ДΤС	ΔΔΔ	CTC	TCC	тст	GTT	GCC	тст	GGA	ጥጥር	ΔΤΤ	ጥጥር	ΔСΤ	AAC	CAC	96
198																	,,
	ser	Met	пуя	Leu	ser	Cys	val	ATA		GTA	FIIE	тте	FILE		ASII	UTR	
199				20					25					30			
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201	TGG	ATG	AAC	TGG	GTC	CGC	CAG	TCT	CCA	GAG	AAG	GGG	CTT	GAG	TGG	GTT	144
202				Trp													
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205																	
200		GAA															192
206 207	Ala	Glu 50	TTE	Arg	ser	ьуs	Ser 55	TIE	Asn	Ser	Ата	Thr 60	His	Tyr	Ala	GIu	
208																	
209		GTG															240
210 211	Ser 65	Val	ьуs	GIY	Arg	Phe 70	Thr	Ile	Ser	Arg	Asp 75	Asp	Ser	Lys	Ser	Ala 80	
212	0.5					70					/5					80	
213	GTC	TAC	CTG	CAA	ATG	ACC	GAC	TTA	AGA	ACT	GAA	GAC	ACT	GGC	GTT	TAT	288
214		Tyr															
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216																	
217 218																	
219	TAC	TGT	TCC	AGG	AAT	TAC	TAC	GGT	AGT	ACC	TAC	GAC	TAC	TGG	GGC	CAA	336
220		Cys															
221	-	-		100		-	-	-	105		-	-	•	110	•		
222																	
223		ACC															357
224 225	Gly	Thr		Leu	Thr	Val	Ser										
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231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247	Ser Trp	(: (x Val Met	ii) M ki) S Lys Lys Asn 35	(A) (B) (B) (D) MOLE(SEQUE Leu 20 Trp	LENCE CULE Solu Solu Solu Val	NGTH: PE: 6 POLOG TYPE DESC Glu Cys Arg	: 119 amino GY: 1 E: pr CRIPT Ser Val	e am: c ac: linea rote: FION Gly Ala Ser 40	ino a id ar in : SE(Gly Ser 25	Q ID Gly 10 Gly Glu	NO:5 Leu Phe Lys	Val Ile Gly	Phe Leu 45	Ser 30 Glu	15 Asn Trp	His Val	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248	Ser Trp	(: (x Val Met	ii) M ki) S Lys Lys Asn 35	(A) (B) (B) (D) MOLE(SEQUE Leu 20 Trp	LENCE CULE Solu Solu Solu Val	NGTH: PE: 6 POLOG TYPE DESC Glu Cys Arg	: 119 amino GY: 1 E: pr CRIPT Ser Val	e am: c ac: linea rote: FION Gly Ala Ser 40	ino a id ar in : SE(Gly Ser 25	Q ID Gly 10 Gly Glu	NO:5 Leu Phe Lys	Val Ile Gly	Phe Leu 45	Ser 30 Glu	15 Asn Trp	His Val	
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231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	Ser Trp	(: (x Val Met Met Glu 50	ii) N ki) S Lys Lys Asn 35	(A) (B) (D) MOLEC SEQUE Leu Leu 20 Trp Arg	LENCE CULE ENCE Glu 5 Ser Val	GTH: Cys Arg	: 119 amino GY: 1 E: pr CRIPT Ser Val Gln Ser 55	e am: c ac: linea rote: rION Gly Ala Ser 40 Ile	ino a id ar in SEG Gly Ser 25 Pro Asn	Gly Gly Glu Ser	NO:5 Leu Phe Lys Ala	Val Ile Gly Thr	Phe Leu 45 His	Ser 30 Glu Tyr	15 Asn Trp Ala	His Val Glu	
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231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	Ser Trp Ala Ser 65	(: (x Val Met Met Glu 50	Lys Lys Asn 35 Ile	(A) (B) (D) MOLEC SEQUI Leu 20 Trp Arg	LENCE CULE ENCE Glu 5 Ser Val Ser	TYPE DESC Glu Cys Arg Lys Phe 70	Ser Ser Thr	e am: c ac: linea rote: FION Gly Ala Ser 40 Ile	ino a id ar in SEG Gly Ser 25 Pro Asn	Cly Clu Ser	NO:5 Leu Phe Lys Ala Asp 75	Val Ile Gly Thr 60 Asp	Phe Leu 45 His	Ser 30 Glu Tyr Lys	15 Asn Trp Ala Ser	His Val Glu Ala 80	

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256 257 258	Tyr	Cys	Ser	Arg 100	Asn	Tyr	Tyr	Gly	Ser 105	Thr	Tyr	Asp	Tyr	Trp 110	Gly	Gln
259 260	Glv	Thr	Thr	T. e 11	Thr	Val	Ser									
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Original Text

SEQUENCE MISSING ITEM REPORT PATENT APPLICATION US/08/013,413A

DATE: 09/17/93 TIME: 15:22:02

INPUT SET: S6568.raw

<< THERE ARE NO ITEMS MISSING >>

SEQUENCE CORRECTION REPORT PATENT APPLICATION US/08/013,413A

DATE: 09/17/93 TIME: 15:22:03

INPUT SET: S6568.raw

Line

Original Text

Corrected Text